

PC-0041 CIP

<110> Yue, Henry
Matthew R. Kaser

<120> PROGESTERONE RECEPTOR COMPLEX P23-LIKE PROTEIN

<130> PC-0041 CIP

<140> To Be Assigned

<141> Herewith

<160> 9

<170> PERL Program

<210> 1

<211> 156

<212> PRT

<213> Homo sapiens

<220>

<221> misc_feature

<223> Incyte ID No: 2923091CD1

<400> 1

Met	Ala	Arg	Gln	His	Ala	Arg	Thr	Leu	Trp	Tyr	Asp	Arg	Pro	Met
1				5					10					15
Tyr	Val	Phe	Met	Glu	Phe	Cys	Val	Glu	Asp	Ser	Thr	Asp	Val	His
			20						25					30
Val	Leu	Ile	Glu	Asp	His	Arg	Ile	Val	Phe	Ser	Cys	Lys	Asn	Ala
			35						40					45
Asp	Gly	Val	Glu	Leu	Tyr	Asn	Glu	Ile	Glu	Phe	Tyr	Ala	Lys	Val
			50						55					60
Asn	Ser	Lys	Asp	Ser	Gln	Asp	Lys	Arg	Ser	Ser	Arg	Ser	Ile	Thr
			65						70					75
Cys	Phe	Val	Arg	Lys	Trp	Lys	Glu	Lys	Val	Ala	Trp	Pro	Arg	Leu
			80						85					90
Thr	Lys	Glu	Asp	Ile	Lys	Pro	Val	Trp	Leu	Ser	Val	Asp	Phe	Asp
			95						100					105
Asn	Trp	Arg	Asp	Trp	Glu	Gly	Asp	Glu	Glu	Met	Glu	Leu	Ala	His
			110						115					120
Val	Glu	His	Tyr	Ala	Glu	Leu	Leu	Lys	Lys	Val	Ser	Thr	Lys	Arg
			125						130					135
Pro	Pro	Pro	Ala	Met	Asp	Asp	Leu	Asp	Asp	Asp	Ser	Asp	Ser	Ala
			140						145					150
Asp	Asp	Ala	Thr	Ser	Asn									
														155

<210> 2

<211> 559

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<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<223> Incyte ID No: 2923091CB1

<400> 2

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ccgcaatggc acggcagcac gcccggaacct tgtggtacga caggcccatg tatgtgttca 60
tggagttttg tgttgaggac agcaccgatg tccacgtgct tattgaggat caccgcattg 120
tggtcagctg caagaatgcc gatggagtgg agttgtacaa tgagattgag ttctatgcca 180
aagtgaactc caaggactcc caggataaagc gctcttcccg ctctattact tgttttgtga 240
gaaaatggaa ggaaaagggtg gcctggccgc ggcttaccaa ggaggatatc aagccagtgt 300
ggctgtctgt ggactttgat aactggagag actgggaagg ggatgaagag atggagctgg 360
ctcatgtgga acattatgca gagcttttga agaaggtcag caccaagaga cctccacctg 420
ccatggatga tttggatgat gattctgaca gtgctgatga tgcaacaagt aattaacttt 480
ctgtgacgca aagctgggaa ggcagctgtg gctattttcc agttgttcta gaaagctagc 540
gcctaggcct ttgtcagcg 559
```

<210> 3

<211> 451

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<223> Incyte ID No: 2184024F6

<400> 3

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ggcacggcag cacgcccga ccttgtggta cgacaggccc atgtatgtgt tcatggagtt 60
ttgtgttgag gacagcaccg atgtccacgt gcttattgag gatcacgca ttgtgttcag 120
ctgcaagaat gccgatggag tggagtgtga caatgagatt gagttctatg ccaaagtga 180
ctccaaggac tcccaggata agcgtcttcc ccgctctatt acttgttttg tgagaaaatg 240
gaaggaaaag gtggcctggc cgcggttac caaggaggat atcaagccag tgtggctgtc 300
tgtggacttt gataactgga gagactggga aggggatgaa gagatggagc tggctcatgt 360
ggaacattat gcagagcttt tgaagaaggt cagcaccaag agacctccac ctgccatgga 420
tgatttggat gatgattctg acagtgtctga t 451
```

<210> 4

<211> 455

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<223> Incyte ID No: 2923091F6

<220>

<221> unsure

<222> 21, 138, 263, 406, 419

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<223> a, t, c, g, or other

<400> 4

```
gcaatggcac ggcagcacgc ncggaccttg tggtaacgaca ggcccaggta tgtgttcattg 60
gagttttgtg ttgaggacag caccgatgtc cacgtgctta ttgaggatca ccgcattgtg 120
ttcagctgca agaatgcnga tggagtggag ttgtacaatg agattgagtt ctatgccaaa 180
gtgaactcca aggactccca ggataagcgc tcttcccgtc ctattacttg ttttgtgaga 240
aaatggaagg aaaagggtggc ctngccgcgg cttaccaagg aggatatcaa gccagtgtgg 300
ctgtctgtgg actttgataa ctggagagac tgggaagggg atgaagagat ggagctggct 360
catgtggaac attatgcaga gcttttgaaa gaaggtcagc accaanagac ctccacctnc 420
catggatgat ttggatggtc tacaagtgtc ttggc 455
```

<210> 5

<211> 231

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<223> Incyte ID No: 3173306H1

<400> 5

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ggaactctaa atgccaccct ggagcgggag ccgcaatggc acggcagcac gcccgacact 60
tgtgtgtacga caggcccagg tatgtgttca tggagttttg tgttgaggac agcaccgatg 120
tccacgtgct tattgaggat caccgcattg tgttcagctg caagaatgcc gatggagtgg 180
agttgtacaa tgagattgag ttctatgcc aagtgaactc caaggactcc c 231
```

<210> 6

<211> 249

<212> DNA

<213> Homo sapiens

<220>

<221> misc_feature

<223> Incyte ID No: 3176831H1

<400> 6

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ggactttgat aactggagag actgggaagg ggatgaagag atggagctgg ctcatgtgga 60
acattatgca gagcttttga agaaggctcag caccaagaga cctccacctg ccatggatga 120
tttgatgat gattctgaca gtgctgatga tgcaacaagt aattaacttt ctgtgacgca 180
aagctgggaa ggcagctgtg gctattttcc agttgttcta gaaagctagc gcctaggcct 240
ttgtcagcg 249
```

<210> 7

<211> 590

<212> DNA

<213> Rattus norvegicus

<220>

<221> misc_feature

PC-0041 CIP

<223> Incyte ID No: 702125891H2

<400> 7

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cctatcacat cgaacctatc actcgagcct atcactcgag ccgatcactc gagcctatca 60
ctcgagcggc agcacgcccg gactctgtgg tacgacagac ccaaataatgt tttcatggag 120
ttttgcgttg aggacagcac tgacgtcagt gtgctcattg aggaccaccg catcgtgttc 180
agctgcagga atggtgatgg tgtggagctg tacaacgaga ttgagttcta tgccaaagtg 240
aactccaagg actcccagga taagcgctct ggctcgctcca ttacttgttt tgtgaggaaa 300
tggaaggaga aggtgccctg gcctcgactc accaaggagg atataaagcc cgtgtggctc 360
tctgtggact tcgataactg gagagactgg gaaggagatg acgagatgga gctggcgag 420
gtggaacact atgcagagct tttgaacaag gtcagcacta agagacctcc ccctgccatg 480
gatgatctgg acgatgattc tgacaactaa ctagctctct gtgacagtgg acctggggag 540
gaggctgtag ctacctctg tcgtgctgag gagctaggat gggctgtcct 590
```

<210> 8

<211> 444

<212> DNA

<213> Mus musculus

<220>

<221> misc_feature

<223> Incyte ID No: 018316_Mm.1

<400> 8

```
gagattgaat tctatgccaa ggtgaactcc aaggactccc aggataagcg ttctgggtcg 60
tccattactt gctttgtgag gaaatggaag gagaagggtg cctggcctag gtcacaaaag 120
gaggatataa agcctgtgtg gctctctgtg gacttcgata actggagaga ctgggaagga 180
gacgatgagg tggagctggc tcaggtggaa cattatgcag agcttctgaa caaggtcagc 240
actaagaggc ctccccctgc catggatgat ctggacgatg attctgacag ctaactagct 300
ttctgtgacg gtggagccgg ggaggaggcg gtacgtatct tctgtcatgc tgaaaaactg 360
ggatgggtgcc ttcttcaact acttggtttg catcaagatc cacagagacc tctgaactct 420
tccagaagct ctttctgaag ggtg 444
```

<210> 9

<211> 160

<212> PRT

<213> Homo sapiens

<220>

<221> misc_feature

<223> Incyte ID No: g438652

<400> 9

```
Met Gln Pro Ala Ser Ala Lys Trp Tyr Asp Arg Arg Asp Tyr Val
  1             5             10             15
Phe Ile Glu Phe Cys Val Glu Asp Ser Lys Asp Val Asn Val Asn
             20             25             30
Phe Glu Lys Ser Lys Leu Thr Phe Ser Cys Leu Gly Gly Ser Asp
             35             40             45
Asn Phe Lys His Leu Asn Glu Ile Asp Leu Phe His Cys Ile Asp
```

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				50					55					60
Pro	Asn	Asp	Ser	Lys	His	Lys	Arg	Thr	Asp	Arg	Ser	Ile	Leu	Cys
				65					70					75
Cys	Leu	Arg	Lys	Gly	Glu	Ser	Gly	Gln	Ser	Trp	Pro	Arg	Leu	Thr
				80					85					90
Lys	Glu	Arg	Ala	Lys	Leu	Asn	Trp	Leu	Ser	Val	Asp	Phe	Asn	Asn
				95					100					105
Trp	Lys	Asp	Trp	Glu	Asp	Asp	Ser	Asp	Glu	Asp	Met	Ser	Asn	Phe
				110					115					120
Asp	Arg	Phe	Ser	Glu	Met	Met	Asn	Asn	Met	Gly	Gly	Asp	Glu	Asp
				125					130					135
Val	Asp	Leu	Pro	Glu	Val	Asp	Gly	Ala	Asp	Asp	Asp	Ser	Gln	Asp
				140					145					150
Ser	Asp	Asp	Glu	Lys	Met	Pro	Asp	Leu	Glu					
				155					160					

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